

appendix_placebo.R

datalab

2023-06-16

```
#####  
#####          Placebo Tests #####  
#####          October 10, 2018 #####  
#####          Rerun December 17, 2022  
  
rm(list=ls())  
library(foreign)  
library(plyr)  
library(readstata13)  
library(multiwayvcov)  
library(sandwich)  
library(lmtest)  
library(stargazer)  
library(ggplot2)  
  
#####  
##### Multiplot Function #####  
##### Source: http://www.cookbook-r.com/Graphs/Multiple\_graphs\_on\_one\_page\_\(ggplot2\)/  
  
multiplot <- function(..., plotlist=NULL, file, cols=1, layout=NULL) {  
  library(grid)  
  
  # Make a list from the ... arguments and plotlist  
  plots <- c(list(...), plotlist)  
  
  numPlots = length(plots)  
  
  # If layout is NULL, then use 'cols' to determine layout  
  if (is.null(layout)) {  
    # Make the panel  
    # ncol: Number of columns of plots  
    # nrow: Number of rows needed, calculated from # of cols  
    layout <- matrix(seq(1, cols * ceiling(numPlots/cols)),  
                      ncol = cols, nrow = ceiling(numPlots/cols))  
  }  
  
  if (numPlots==1) {  
    print(plots[[1]])  
  }  
}
```

```

} else {
  # Set up the page
  grid.newpage()
  pushViewport(viewport(layout = grid.layout(nrow(layout), ncol(layout))))

  # Make each plot, in the correct location
  for (i in 1:numPlots) {
    # Get the i,j matrix positions of the regions that contain this subplot
    matchidx <- as.data.frame(which(layout == i, arr.ind = TRUE))

    print(plots[[i]], vp = viewport(layout.pos.row = matchidx$row,
                                     layout.pos.col = matchidx$col))
  }
}
}
#####

data=read.csv("~/Dropbox/Personal Research 2017/replications/karn_nov16.csv")
names(data)

```

```

## [1] "X.1"           "dist_name"      "vilname91"      "v1"             "dist_code"
## [15] "phc_cntr"       "hc_cntr"        "fpc_cntr"       "tb_cntr"        "nh_cntr"
## [29] "st_town"        "agri_land"      "near_town"      "circl_code"     "m_pop"
## [43] "m_sc"           "f_sc"           "tot_st"         "m_st"           "f_st"
## [57] "ngmf_char"      "ngmfprwmedd"    "ngmfprwothd"    "ngmfprwnod"     "ngmftrpr"
## [71] "taptr"          "tapuntr"        "hp"             "cowwell"        "uncovwell"
## [85] "phs_cntr"       "prhsc"          "stname"         "stname1991"     "d_name"
## [99] "all_hosp"       "area_na_cu"     "ayu_disp"       "ayu_hosp"       "canal_govt"
## [113] "ind_sch"        "lake"           "m_home"         "m_sch"          "nw_fac"
## [127] "power_oth"      "power_supl"     "p_sch"          "p_t_fac"        "rang_mcw"
## [141] "s_sch"          "s_s_sch"        "tot_exp"        "tot_inc"        "tr_sch"
## [155] "gov_ps_n"       "pr_ps_n"        "gov_ms_n"       "pr_ms_n"        "gov_secs_n"
## [169] "nviltmsna"      "nviltsecs"     "nviltsecsna"   "power"          "hplost"
## [183] "pnt_fac"        "power_ea"       "power_eag"     "power_edea"     "power_eo"
## [197] "medfac"         "rangmed"        "tot_hh"         "pucca_binary"   "kucha_binary"
## [211] "dist_fr_town"   "tbcl"           "tank"           "tap"            "X"
## [225] "M_POP"          "F_POP"          "TOT_L6"         "M_L6"           "F_L6"
## [239] "TOT_ILLT"       "M_ILLT"         "F_ILLT"         "TOT_W"          "M_W"
## [253] "F_AGLB"         "TOT_MFHH"       "M_MFHH"         "F_MFHH"         "TOT_OTH_W"
## [267] "M_MRG_AGLB"     "F_MRG_AGLB"    "T_MRG_HH"       "M_MRG_HH"       "F_MRG_HH"
## [281] "NEAR_DIST_border1" "NEAR_ANGLE"    "temp_av"        "wc2010mt_1"    "TerrainRug"

```

```
summary(data$NEAR_DIST_border1)
```

```

##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.    NA's
## 45.1  17162.2 40186.1 45728.7 67400.2 133049.0 5146

```

```
summary(data$border1)
```

```

##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.    NA's
## 0.000  0.000   1.000  0.599  1.000   1.000  5146

```

```

#####
###Create Placebo Border to the North (-10 km to Princely State)

```

```
#calculate new distances
```

```
#border1
```

```
#mysore part
```

```
summary(data$NEAR_DIST_border1[data$border1==1])
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.   NA's  
##  119.9 20224.4 42551.1 47531.3 66465.8 133049.0 5146
```

```
data$distplacebo1[(data$border1==1)&!is.na(data$border1)]=  
  data$NEAR_DIST_border1[(data$border1==1)&!is.na(data$border1)]+10000
```

```
summary(data$distplacebo1[data$border1==1])
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.   NA's  
##  10120 30224 52551 57531 76466 143049 5146
```

```
#bombay part
```

```
data$distplacebo1[(data$border1==0)&!is.na(data$border1)  
  &(data$NEAR_DIST_border1<10000)]=  
  10000-data$NEAR_DIST_border1[(data$border1==0)&!is.na(data$border1)  
  &(data$NEAR_DIST_border1<10000)]
```

```
data$distplacebo1[(data$border1==0)&!is.na(data$border1)  
  &(data$NEAR_DIST_border1>=10000)]=  
  data$NEAR_DIST_border1[(data$border1==0)&!is.na(data$border1)  
  &(data$NEAR_DIST_border1>=10000)]-10000
```

```
summary(data$NEAR_DIST_border1[data$border1==0])
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.   NA's  
##  45.1 13709.4 34703.0 43034.2 68323.9 130614.8 5146
```

```
summary(data$distplacebo1[data$border1==0])
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.   NA's  
##  3.57 7355.51 24702.97 34974.50 58323.93 120614.78 5146
```

```
#create a new treatment variable
```

```
#old mysore and chunk of the bombay are in the treatment,  
#and the rest of bombay are in teh control group
```

```
data$borderplac1[data$border1==1]=1  
data$borderplac1[(data$border1==0)&!is.na(data$border1)  
  &(data$NEAR_DIST_border1<10000)]=1  
data$borderplac1[(data$border1==0)&!is.na(data$border1)  
  &(data$NEAR_DIST_border1>=10000)]=0
```

```
summary(data$borderplac1)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.   NA's  
##  0.000 0.000 1.000 0.674 1.000 1.000 5146
```

```
##### Distances #####
```

```
#Distance to Mysore-Bombay Border
```

```
rd10.mb=data[which(data$distplacebo1<10000),] #20 km
```

```
table(rd10.mb$borderplac1)
```

```

##
## 0 1
## 455 559
#####Linear Polynomial #####

#baseline bandwidth
#Mysore-Bombay
mys.health1=lm(health_binary~borderplac1+TOT_POP+
              TOT_SC+TOT_ST+Slope+TerrainRug+Latitude+Longitude, data=rd10.mb)
summary(mys.health1)

##
## Call:
## lm(formula = health_binary ~ borderplac1 + TOT_POP + TOT_SC +
##     TOT_ST + Slope + TerrainRug + Latitude + Longitude, data = rd10.mb)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.7468 -0.2738 -0.1436  0.3224  1.0358
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -8.571e+00  3.430e+00  -2.499  0.01261 *
## borderplac1 -5.106e-02  2.823e-02  -1.809  0.07075 .
## TOT_POP      9.626e-05  1.383e-05   6.962 6.04e-12 ***
## TOT_SC       1.272e-04  4.370e-05   2.911  0.00368 **
## TOT_ST       2.871e-04  5.617e-05   5.111 3.84e-07 ***
## Slope       -2.702e-04  1.159e-03  -0.233  0.81569
## TerrainRug  -2.827e-03  6.345e-03  -0.446  0.65598
## Latitude    -1.787e-01  7.933e-02  -2.253  0.02448 *
## Longitude    1.499e-01  5.146e-02   2.912  0.00367 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.4083 on 1005 degrees of freedom
## Multiple R-squared:  0.1399, Adjusted R-squared:  0.133
## F-statistic: 20.43 on 8 and 1005 DF,  p-value: < 2.2e-16

mys.health1.cl=cluster.vcov(mys.health1, rd10.mb$dist_name)
mys.health1.se=sqrt(diag(mys.health1.cl)) #cluster standard errors

mys.pucca1=lm(pucca_binary~borderplac1+TOT_POP+
             TOT_SC+TOT_ST+Slope+TerrainRug+Latitude+Longitude, data=rd10.mb)
summary(mys.pucca1)

##
## Call:
## lm(formula = pucca_binary ~ borderplac1 + TOT_POP + TOT_SC +
##     TOT_ST + Slope + TerrainRug + Latitude + Longitude, data = rd10.mb)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.00046  0.01794  0.08377  0.13515  0.23091
##

```

```

## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept) -8.057e+00  2.355e+00  -3.422 0.000647 ***
## borderplac1  9.152e-04  1.938e-02   0.047 0.962341
## TOT_POP      1.550e-05  9.492e-06   1.633 0.102829
## TOT_SC       2.281e-05  3.000e-05   0.760 0.447169
## TOT_ST       5.406e-05  3.856e-05   1.402 0.161287
## Slope        1.551e-04  7.955e-04   0.195 0.845441
## TerrainRug   -4.616e-03  4.356e-03  -1.060 0.289529
## Latitude     -4.469e-02  5.447e-02  -0.820 0.412142
## Longitude    1.275e-01  3.533e-02   3.610 0.000322 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.2803 on 1005 degrees of freedom
## Multiple R-squared:  0.04659,    Adjusted R-squared:  0.039
## F-statistic: 6.139 on 8 and 1005 DF,  p-value: 9.476e-08

```

```

mys.pucca1.cl=cluster.vcov(mys.pucca1, rd10.mb$dist_name)
mys.pucca1.se=sqrt(diag(mys.pucca1.cl)) #cluster standard errors

```

```

#####
#### Create Placebo Border to the South (+10 km to the princely state)

```

```

#border1
#mysore part
summary(data$NEAR_DIST_border1[data$border1==0])

```

```

##      Min.   1st Qu.   Median     Mean  3rd Qu.     Max.    NA's
##      45.1  13709.4  34703.0  43034.2  68323.9 130614.8  5146

```

```

data$distplacebo1[(data$border1==0)&!is.na(data$border1)]=
  data$NEAR_DIST_border1[(data$border1==0)&!is.na(data$border1)]+10000

```

```

summary(data$distplacebo1[data$border1==0])

```

```

##      Min.   1st Qu.   Median     Mean  3rd Qu.     Max.    NA's
##      10045  23709   44703   53034   78324   140615   5146

```

```

#bombay part
data$distplacebo1[(data$border1==1)&!is.na(data$border1)
  &(data$NEAR_DIST_border1<10000)]=
  10000-data$NEAR_DIST_border1[(data$border1==1)&!is.na(data$border1)
  &(data$NEAR_DIST_border1<10000)]

```

```

data$distplacebo1[(data$border1==1)&!is.na(data$border1)
  &(data$NEAR_DIST_border1>=10000)]=
  data$NEAR_DIST_border1[(data$border1==1)&!is.na(data$border1)
  &(data$NEAR_DIST_border1>=10000)]-10000

```

```

summary(data$NEAR_DIST_border1[data$border1==1])

```

```

##      Min.   1st Qu.   Median     Mean  3rd Qu.     Max.    NA's
##      119.9  20224.4  42551.1  47531.3  66465.8 133049.0  5146

```

```

summary(data$distplacebo1[data$border1==1])

##      Min.   1st Qu.   Median     Mean   3rd Qu.     Max.      NA's
##    12.96  10224.44  32551.09 38913.18 56465.82 123049.02 5146

#create a new treatment variable
#old mysore and chunk of the bombay are in the treatment,
#and the rest of bombay are in teh control group

data$borderplac1[data$border1==0]=1
data$borderplac1[(data$border1==1)&!is.na(data$border1)
                 &(data$NEAR_DIST_border1<10000)]=1
data$borderplac1[(data$border1==1)&!is.na(data$border1)
                 &(data$NEAR_DIST_border1>=10000)]=0
summary(data$borderplac1)

##      Min. 1st Qu.  Median     Mean 3rd Qu.     Max.      NA's
##    0.000  0.000  0.000  0.481  1.000  1.000  5146

##### Distances #####

#Distance to Mysore-Bombay Border
rd10.mb=data[which(data$distplacebo1<10000),] #20 km

table(rd10.mb$borderplac1)

##
##    0  1
## 511 599

##### Simple OLS with controls #####
## and linear polynomial of lat and long ###

#baseline bandwidth
#Mysore-Bombay
mys.health2=lm(health_binary~borderplac1+TOT_POP+
               TOT_SC+TOT_ST+Slope+TerrainRug+Latitude+Longitude, data=rd10.mb)
summary(mys.health2)

##
## Call:
## lm(formula = health_binary ~ borderplac1 + TOT_POP + TOT_SC +
##     TOT_ST + Slope + TerrainRug + Latitude + Longitude, data = rd10.mb)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.52151 -0.23538 -0.13628 -0.05932  1.00326
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -9.155e+00  3.085e+00  -2.968  0.00307 **
## borderplac1  6.455e-03  2.722e-02   0.237  0.81262
## TOT_POP     -2.105e-06  2.080e-05  -0.101  0.91943
## TOT_SC       1.824e-04  4.725e-05   3.861  0.00012 ***
## TOT_ST       3.298e-04  6.071e-05   5.432  6.84e-08 ***

```

```

## Slope      1.701e-03  1.059e-03  1.607  0.10843
## TerrainRug 8.472e-04  7.005e-03  0.121  0.90376
## Latitude   1.525e-01  8.493e-02  1.795  0.07286 .
## Longitude  9.180e-02  4.225e-02  2.173  0.03001 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3929 on 1101 degrees of freedom
## Multiple R-squared:  0.07367,    Adjusted R-squared:  0.06694
## F-statistic: 10.95 on 8 and 1101 DF,  p-value: 6.085e-15

mys.health2.cl=cluster.vcov(mys.health2, rd10.mb$dist_name)
mys.health2.se=sqrt(diag(mys.health2.cl)) #cluster standard errors

mys.pucca2=lm(pucca_binary~borderplac1+TOT_POP+
             TOT_SC+TOT_ST+Slope+TerrainRug+Latitude+Longitude, data=rd10.mb)
summary(mys.pucca2)

##
## Call:
## lm(formula = pucca_binary ~ borderplac1 + TOT_POP + TOT_SC +
##     TOT_ST + Slope + TerrainRug + Latitude + Longitude, data = rd10.mb)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.03623 -0.01632  0.16068  0.25919  0.52007
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.479e-01  3.157e+00  -0.047  0.962656
## borderplac1  3.281e-02  2.786e-02   1.178  0.239244
## TOT_POP      1.322e-04  2.129e-05   6.212  7.41e-10 ***
## TOT_SC       2.330e-04  4.836e-05   4.817  1.66e-06 ***
## TOT_ST       2.322e-04  6.213e-05   3.738  0.000195 ***
## Slope        3.137e-03  1.084e-03   2.894  0.003880 **
## TerrainRug   -7.225e-03  7.170e-03  -1.008  0.313837
## Latitude     1.606e-02  8.692e-02   0.185  0.853428
## Longitude    2.421e-03  4.324e-02   0.056  0.955369
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.4022 on 1101 degrees of freedom
## Multiple R-squared:  0.09589,    Adjusted R-squared:  0.08932
## F-statistic: 14.6 on 8 and 1101 DF,  p-value: < 2.2e-16

mys.pucca2.cl=cluster.vcov(mys.pucca2, rd10.mb$dist_name)
mys.pucca2.se=sqrt(diag(mys.pucca2.cl)) #cluster standard errors

#####
#border2

####Calculate new distances (-10 km from the princely state)

```

```

#hyd part
summary(data$NEAR_DIST_border2[data$border2==1])

##      Min.   1st Qu.   Median     Mean  3rd Qu.    Max.     NA's
## 307.7  21747.8  46276.5  49566.0  73561.2 139213.6  6425

data$distplacebo2[(data$border2==1)&!is.na(data$border2)]=
  data$NEAR_DIST_border2[(data$border2==1)&!is.na(data$border2)]+10000

summary(data$distplacebo2[data$border2==1])

##      Min.   1st Qu.   Median     Mean  3rd Qu.    Max.     NA's
## 10308  31748   56276   59566   83561   149214   6425

#bombay part
data$distplacebo2[(data$border2==0)&!is.na(data$border2)
  &(data$NEAR_DIST_border2<10000)]=
  10000-data$NEAR_DIST_border2[(data$border2==0)&!is.na(data$border2)
  &(data$NEAR_DIST_border2<10000)]

data$distplacebo2[(data$border2==0)&!is.na(data$border2)
  &(data$NEAR_DIST_border2>=10000)]=
  data$NEAR_DIST_border2[(data$border2==0)&!is.na(data$border2)
  &(data$NEAR_DIST_border2>=10000)]-10000

summary(data$NEAR_DIST_border2[data$border2==0])

##      Min.   1st Qu.   Median     Mean  3rd Qu.    Max.     NA's
## 235.7  17397.5  40491.5  45437.5  74356.1 111267.2  6425

summary(data$distplacebo2[data$border2==0])

##      Min.   1st Qu.   Median     Mean  3rd Qu.    Max.     NA's
## 19.45  8982.04  30491.49  37324.40  64356.14 101267.22  6425

#create a new treatment variable
#old mysore and chunk of the bombay are in the treatment,
#and the rest of bombay are in teh control group
data$borderplac2[data$border2==1]=1
data$borderplac2[(data$border2==0)&!is.na(data$border2)
  &(data$NEAR_DIST_border2<10000)]=1
data$borderplac2[(data$border2==0)&!is.na(data$border2)
  &(data$NEAR_DIST_border2>=10000)]=0
summary(data$borderplac2)

##      Min.   1st Qu.   Median     Mean  3rd Qu.    Max.     NA's
## 0.000  0.000   1.000   0.641   1.000   1.000   6425

#Distance to Hyderabad-Bombay Border
rd10.hb=data[which(data$distplacebo2<10000),] #20 km
table(rd10.hb$borderplac2)

##
## 0 1
## 293 447

#hyderabad

```

```

hyd.health1=lm(health_binary~borderplac2+TOT_POP+
              TOT_SC+TOT_ST+Slope+TerrainRug+Latitude+Longitude, data=rd10.hb)
summary(hyd.health1)

```

```

##
## Call:
## lm(formula = health_binary ~ borderplac2 + TOT_POP + TOT_SC +
##     TOT_ST + Slope + TerrainRug + Latitude + Longitude, data = rd10.hb)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.7378 -0.3469 -0.2292  0.4792  0.9156
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -2.010e+00  7.881e+00  -0.255  0.7988
## borderplac2 -9.133e-03  3.814e-02  -0.239  0.8108
## TOT_POP      1.109e-04  1.715e-05   6.468 1.82e-10 ***
## TOT_SC       1.088e-04  5.169e-05   2.104  0.0357 *
## TOT_ST       1.838e-04  7.360e-05   2.498  0.0127 *
## Slope        -4.688e-04  1.006e-03  -0.466  0.6413
## TerrainRug   -1.031e-02  2.011e-02  -0.513  0.6083
## Latitude     -2.376e-02  4.253e-02  -0.559  0.5765
## Longitude     3.331e-02  1.095e-01   0.304  0.7611
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.4579 on 731 degrees of freedom
## Multiple R-squared:  0.08649,    Adjusted R-squared:  0.07649
## F-statistic: 8.651 on 8 and 731 DF,  p-value: 2.558e-11

```

```

hyd.health1.cl=cluster.vcov(hyd.health1, rd10.hb$dist_name)
hyd.health1.se=sqrt(diag(hyd.health1.cl)) #cluster standard errors

```

```

hyd.pucca1=lm(pucca_binary~borderplac2+TOT_POP+
              TOT_SC+TOT_ST+Slope+TerrainRug+Latitude+Longitude, data=rd10.hb)
summary(hyd.pucca1)

```

```

##
## Call:
## lm(formula = pucca_binary ~ borderplac2 + TOT_POP + TOT_SC +
##     TOT_ST + Slope + TerrainRug + Latitude + Longitude, data = rd10.hb)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.01086  0.02143  0.08926  0.13991  0.25204
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  8.087e+00  5.076e+00   1.593 0.111541
## borderplac2 -1.297e-02  2.456e-02  -0.528 0.597714
## TOT_POP      4.268e-05  1.105e-05   3.864 0.000122 ***
## TOT_SC       4.674e-05  3.329e-05   1.404 0.160797
## TOT_ST       8.877e-05  4.740e-05   1.873 0.061501 .

```

```

## Slope      -4.650e-04  6.479e-04  -0.718  0.473208
## TerrainRug  2.770e-02  1.295e-02   2.139  0.032764 *
## Latitude   -1.608e-02  2.739e-02  -0.587  0.557391
## Longitude  -9.228e-02  7.053e-02  -1.308  0.191149
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.2949 on 731 degrees of freedom
## Multiple R-squared:  0.05689,    Adjusted R-squared:  0.04656
## F-statistic: 5.511 on 8 and 731 DF,  p-value: 8.832e-07
hyd.pucca1.cl=cluster.vcov(hyd.pucca1, rd10.hb$dist_name)
hyd.pucca1.se=sqrt(diag(hyd.pucca1.cl)) #cluster standard errors

###Calculate new distances (+10 km to the princely state)

#hyd part
summary(data$NEAR_DIST_border2[data$border2==0])

##      Min.   1st Qu.   Median     Mean   3rd Qu.     Max.    NA's
##   235.7  17397.5  40491.5  45437.5  74356.1 111267.2  6425
data$distplacebo2[(data$border2==0)&!is.na(data$border2)]=
  data$NEAR_DIST_border2[(data$border2==0)&!is.na(data$border2)]+10000

summary(data$distplacebo2[data$border2==0])

##      Min.   1st Qu.   Median     Mean   3rd Qu.     Max.    NA's
##   10236   27397   50491   55437   84356   121267   6425

#bombay part
data$distplacebo2[(data$border2==1)&!is.na(data$border2)
  &(data$NEAR_DIST_border2<10000)]=
  10000-data$NEAR_DIST_border2[(data$border2==1)&!is.na(data$border2)
  &(data$NEAR_DIST_border2<10000)]

data$distplacebo2[(data$border2==1)&!is.na(data$border2)
  &(data$NEAR_DIST_border2>=10000)]=
  data$NEAR_DIST_border2[(data$border2==1)&!is.na(data$border2)
  &(data$NEAR_DIST_border2>=10000)]-10000

summary(data$NEAR_DIST_border2[data$border2==1])

##      Min.   1st Qu.   Median     Mean   3rd Qu.     Max.    NA's
##   307.7  21747.8  46276.5  49566.0  73561.2 139213.6  6425

summary(data$distplacebo2[data$border2==1])

##      Min.   1st Qu.   Median     Mean   3rd Qu.     Max.    NA's
##      0.82  11747.78  36276.47  41199.57  63561.22 129213.56  6425

#create a new treatment variable
#old mysore and chunk of the bombay are in the treatment,
#and the rest of bombay are in teh control group
data$borderplac2[data$border2==0]=1
data$borderplac2[(data$border2==1)&!is.na(data$border2)]

```

```

      &(data$NEAR_DIST_border2<10000)]=1
data$borderplac2[(data$border2==1)&!is.na(data$border2)
      &(data$NEAR_DIST_border2>=10000)]=0
summary(data$borderplac2)

##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.    NA's
##      0.000  0.000   1.000   0.511   1.000   1.000   6425

#Distance to Hyderabad-Bombay Border
rd10.hb=data[which(data$distplacebo2<10000),] #20 km
table(rd10.hb$borderplac2)

##
##      0      1
## 326 493

#hyderabad
hyd.health2=lm(health_binary~borderplac2+TOT_POP+
      TOT_SC+TOT_ST+Slope+TerrainRug+Latitude+Longitude, data=rd10.hb)
summary(hyd.health2)

##
## Call:
## lm(formula = health_binary ~ borderplac2 + TOT_POP + TOT_SC +
##     TOT_ST + Slope + TerrainRug + Latitude + Longitude, data = rd10.hb)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.60432 -0.26117 -0.15722  0.03745  0.98974
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  2.306e+01  7.321e+00   3.150 0.001691 **
## borderplac2  -5.162e-02  3.272e-02  -1.577 0.115120
## TOT_POP      5.248e-05  1.727e-05   3.038 0.002457 **
## TOT_SC       3.319e-04  4.763e-05   6.970 6.59e-12 ***
## TOT_ST       1.950e-04  5.476e-05   3.561 0.000391 ***
## Slope        1.236e-04  1.096e-03   0.113 0.910253
## TerrainRug   1.594e-02  1.574e-02   1.013 0.311505
## Latitude     1.281e-01  3.805e-02   3.368 0.000793 ***
## Longitude   -3.296e-01  1.017e-01  -3.240 0.001244 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3985 on 810 degrees of freedom
## Multiple R-squared:  0.1119, Adjusted R-squared:  0.1031
## F-statistic: 12.76 on 8 and 810 DF,  p-value: < 2.2e-16

hyd.health2.cl=cluster.vcov(hyd.health2, rd10.hb$dist_name)
hyd.health2.se=sqrt(diag(hyd.health2.cl)) #cluster standard errors

hyd.pucca2=lm(pucca_binary~borderplac2+TOT_POP+
      TOT_SC+TOT_ST+Slope+TerrainRug+Latitude+Longitude, data=rd10.hb)
summary(hyd.pucca2)

##

```

```

## Call:
## lm(formula = pucca_binary ~ borderplac2 + TOT_POP + TOT_SC +
##     TOT_ST + Slope + TerrainRug + Latitude + Longitude, data = rd10.hb)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.02943  0.03998  0.10971  0.16980  0.39534
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  1.711e+00  6.080e+00   0.281  0.77850
## borderplac2  1.015e-02  2.718e-02   0.374  0.70886
## TOT_POP      6.068e-05  1.435e-05   4.229 2.61e-05 ***
## TOT_SC      1.167e-04  3.956e-05   2.949  0.00328 **
## TOT_ST      7.076e-05  4.548e-05   1.556  0.12012
## Slope       -1.364e-03  9.106e-04  -1.498  0.13452
## TerrainRug  -1.404e-02  1.307e-02  -1.074  0.28310
## Latitude    6.979e-02  3.160e-02   2.209  0.02749 *
## Longitude   -2.608e-02  8.450e-02  -0.309  0.75768
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.331 on 810 degrees of freedom
## Multiple R-squared:  0.0609, Adjusted R-squared:  0.05162
## F-statistic: 6.566 on 8 and 810 DF,  p-value: 2.534e-08
hyd.pucca2.cl=cluster.vcov(hyd.pucca2, rd10.hb$dist_name)
hyd.pucca2.se=sqrt(diag(hyd.pucca2.cl)) #cluster standard errors

stargazer(mys.health1, mys.pucca1, mys.health2, mys.pucca2,
          se=list(mys.health1.se, mys.pucca1.se, mys.health2.se, mys.pucca2.se), digits=3,
          omit=c("TOT_POP", "TOT_SC", "TOT_ST", "Slope", "TerrainRug", "Latitude", "Longitude"),
          dep.var.labels=c("Health Centers", "Paved Roads", "Health Centers", "Paved Roads"),
          column.labels =c("-10 km", "+10 km"), column.separate = c(2,2),
          covariate.labels = c("Placebo Indirect Rule", "Constant"),
          add.lines = list(c("Controls", "\\checkbox", "\\checkbox", "\\checkbox", "\\checkbox")),
          omit.stat = c("rsq", "f", "adj.rsq", "ser"))

##
## % Table created by stargazer v.5.2.2 by Marek Hlavac, Harvard University. E-mail: hlavac at fas.harvard.edu
## % Date and time: Fri, Jun 16, 2023 - 15:10:45
## \begin{table}[!htbp] \centering
##   \caption{}
##   \label{}
##   \begin{tabular}{@{\extracolsep{5pt}}lcccc}
##     \hline
##     \hline \hline \hline
##     & \multicolumn{4}{c}{\textit{Dependent variable:}} & \\
##     \cline{2-5}
##     \hline & Health Centers & Paved Roads & Health Centers & Paved Roads & \\
##     & \multicolumn{2}{c}{-10 km} & \multicolumn{2}{c}{+10 km} & \\
##     \hline & (1) & (2) & (3) & (4) & \\

```

```
## \hline \[-1.8ex]
## Placebo Indirect Rule & $-0.051 & 0.001 & 0.006 & 0.033 \\  
## & (0.032) & (0.027) & (0.007) & (0.021) \\  
## & & & & \\  
## Constant & $-8.571$^{***}$ & $-8.057$^{***}$ & $-9.155 & $-0.148 \\  
## & (0.474) & (2.923) & (5.599) & (0.970) \\  
## & & & & \\  
## \hline \[-1.8ex]
## Controls & \checkmark & \checkmark & \checkmark & \checkmark \\  
## Observations & 1,014 & 1,014 & 1,110 & 1,110 \\  
## \hline
## \hline \[-1.8ex]
## \textit{Note:} & \multicolumn{4}{r}{\textsuperscript{*}$p$<$0.1; \textsuperscript{**}$p$<$0.05; \textsuperscript{***}$p$<$0.01} \\  
## \end{tabular}
## \end{table}
```

```
stargazer(hyd.health1, hyd.pucca1, hyd.health2, hyd.pucca2,  
          se=list(hyd.health1.se, hyd.pucca1.se, hyd.health2.se, hyd.pucca2.se), digits=3,  
          omit=c("TOT_POP", "TOT_SC", "TOT_ST", "Slope", "TerrainRug", "Latitude", "Longitude"),  
          dep.var.labels=c("Health Centers", "Paved Roads", "Health Centers", "Paved Roads"),  
          column.labels =c("-10 km", "+10 km"), column.separate = c(2,2),  
          covariate.labels = c("Placebo Indirect Rule", "Constant"),  
          add.lines = list(c("Controls", "\\checkmark", "\\checkmark", "\\checkmark", "\\checkmark")),  
          omit.stat = c("rsq", "f", "adj.rsq", "ser"))
```

```
##  
## % Table created by stargazer v.5.2.2 by Marek Hlavac, Harvard University. E-mail: hlavac at fas.harvard.edu  
## % Date and time: Fri, Jun 16, 2023 - 15:10:45  
## \begin{table}[!htbp] \centering  
## \caption{  
## \label{  
## \begin{tabular}{@{\extracolsep{5pt}}lcccc}  
## \[-1.8ex]\hline  
## \hline \[-1.8ex]  
## & \multicolumn{4}{c}{\textit{Dependent variable:}} \\  
## \cline{2-5}  
## \[-1.8ex] & Health Centers & Paved Roads & Health Centers & Paved Roads \\  
## & \multicolumn{2}{c}{-10 km} & \multicolumn{2}{c}{+10 km} \\  
## \[-1.8ex] & (1) & (2) & (3) & (4) \\  
## \hline \[-1.8ex]  
## Placebo Indirect Rule & $-0.009 & $-0.013 & $-0.052$^{***}$ & 0.010 \\  
## & (0.041) & (0.028) & (0.011) & (0.020) \\  
## & & & & \\  
## Constant & $-2.010 & 8.087$^{***}$ & 23.063$^{**}$ & 1.711 \\  
## & (3.585) & (1.776) & (10.229) & (8.803) \\  
## & & & & \\  
## \hline \[-1.8ex]  
## Controls & \checkmark & \checkmark & \checkmark & \checkmark \\  
## Observations & 740 & 740 & 819 & 819 \\  
## \hline  
## \hline \[-1.8ex]  
## \textit{Note:} & \multicolumn{4}{r}{\textsuperscript{*}$p$<$0.1; \textsuperscript{**}$p$<$0.05; \textsuperscript{***}$p$<$0.01} \\  
## \end{tabular}  
## \end{table}
```